

GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: May 4, 2005, 12:02:42 ; Search time 455.392  
Seconds

(without alignments)  
1814.018 Million cell

updates/sec

Title: US-09-300-482-14  
Perfect score: 410  
Sequence: 1  
cccacgcgtccgcggtcatg.....gcaacggcccgctccttgatg 410

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 9369276 seqs, 1007425964 residues

Total number of hits satisfying chosen parameters: 18738552

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Pending\_Patents\_NA\_New:\*  
1: /cgn2\_6/ptodata/1/pna/PCT\_NEW\_COMB.seq:\*  
2: /cgn2\_6/ptodata/1/pna/PCT\_NEW\_COMB.seq2:\*  
3: /cgn2\_6/ptodata/1/pna/US06\_NEW\_COMB.seq:\*  
4: /cgn2\_6/ptodata/1/pna/US07\_NEW\_COMB.seq:\*  
5: /cgn2\_6/ptodata/1/pna/US08\_NEW\_COMB.seq:\*  
6: /cgn2\_6/ptodata/1/pna/US09\_NEW\_COMB.seq:\*  
7: /cgn2\_6/ptodata/1/pna/US10\_NEW\_COMB.seq:\*  
8: /cgn2\_6/ptodata/1/pna/US10\_NEW\_COMB.seq2:\*  
9: /cgn2\_6/ptodata/1/pna/US10\_NEW\_COMB.seq3:\*  
10: /cgn2\_6/ptodata/1/pna/US11\_NEW\_COMB.seq:\*  
11: /cgn2\_6/ptodata/1/pna/US11\_NEW\_COMB.seq2:\*  
12: /cgn2\_6/ptodata/1/pna/US11\_NEW\_COMB.seq3:\*  
13: /cgn2\_6/ptodata/1/pna/US60\_NEW\_COMB.seq:\*